# This Page Is Inserted by IFW Operations and is not a part of the Official Record

## **BEST AVAILABLE IMAGES**

Defective images within this document are accurate representations of the original documents submitted by the applicant.

Defects in the images may include (but are not limited to):

- BLACK BORDERS
- TEXT CUT OFF AT TOP, BOTTOM OR SIDES
- FADED TEXT
- ILLEGIBLE TEXT
- SKEWED/SLANTED IMAGES
- COLORED PHOTOS
- BLACK OR VERY BLACK AND WHITE DARK PHOTOS
- GRAY SCALE DOCUMENTS

## IMAGES ARE BEST AVAILABLE COPY.

As rescanning documents will not correct images, please do not report the images to the Image Problem Mailbox.



### (12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

## (19) World Intellectual Property Organization International Bureau



## 

## (43) International Publication Date 22 March 2001 (22.03.2001)

**PCT** 

## (10) International Publication Number WO 01/1998 A1

- (51) International Patent Classification<sup>7</sup>: C12N 15/31, 15/09, 1/21, A61K 39/108 // (C12N 1/21, C12R 1:19)
- (21) International Application Number: PCT/KR99/00555
- (22) International Filing Date:

15 September 1999 (15.09.1999)

(25) Filing Language:

English

(26) Publication Language:

English

- (71) Applicant (for all designated States except US): MOGAM BIOTECHNOLOGY RESEARCH INSTITUTE [KR/KR]; 341, Pojung-ri, Koosung-myon, Yongin-si, Kyonggi-do 449-910 (KR).
- (72) Inventors; and
- (75) Inventors/Applicants (for US only): PARK, Eun, Jeong [KR/KR]; 322-1007, Woosung Apt., Sangrok-maeul, Jungja-dong, Pundang-ku, Sungnam-si, Kyonggi-do 464-010 (KR). KIM, Jang, Seong [KR/KR]; 902 Ikjoo Apt., Yeonmoo-dong, Jangan-ku, Suwon, Kyonggi-do 440-240 (KR). CHANG, Jihoon [KR/KR]; 553-517, Jayang 3-dong, Kwangjin-ku, Seoul 143-193 (KR). YUM, Jungsun [KR/KR]; 115-502, Chunggu Apt., Hansol-maeul, Jungja-dong, Pundang-ku, Sungnam-si,

Kyonggi-di 463-010 (KR). CHUNG, Soo-il [US/KR]; 112-902, Hyundae Apt., Hyoja-chon, Seohyun-dong, Pundang-ku, Sungnam-si, Kyonggi-do 463-480 (KR).

- (74) Agent: LEE, Han-Young; Seowon Building, 8th floor, 1675-1, Seocho-dong, Seocho-gu, Seoul 137-070 (KR).
- (81) Designated States (national): AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, DE, DK, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, UA, UG, US, UZ, VN, YU, ZA, ZW.
- (84) Designated States (regional): ARIPO patent (GH, GM, KE, LS, MW, SD, SL, SZ, TZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG).

### Published:

With international search report.

For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

-- (5

(54) Title: NOVEL DETOXIFIED MUTANTS OF ESCHERICHIA COLI HEAT-LABILE ENTEROTOXIN

(57) Abstract: The present invention relates to detoxified and immunologically active proteins ("mutant LTs") having mutated amino acid sequences of heat-labile enterotoxin of *E. coli*, DNA sequences encoding the mutant LTs, recombinant expression vectors comprising the DNAs, recombinant microorganisms transformed with the recombinant expression vectors, process for preparing the mutant LTs and pharmaceutical application of the said protein as immunogenic antigens for vaccination and as adjuvants for antibody production. In contrast to wild-type LT, the mutant LTs did not induce any toxic activities. The mutant LTs elicited high and comparable levels of anti-LT antibodies when delivered either intragastrically or intranasally, inducing systemic and local responses in serum and fecal extracts. Thus, they might be useful for the development of a novel diarrheal vaccine in humans and animals. In addition, the antibody production ability using mutant LTs as an adjuvant may be effective for prevention and treatment of various diseases.





### NOVEL DETOXIFITED MUTANTS OF Escherichia coli HEAT-LABILE ENTEROTOXIN

### BACKGROUND OF THE INVENTION

### Field of the Invention

immunologically active proteins ("mutant LTs"), more specifically, to mutant LTs having mutated amino acid sequences of heat-labile enterotoxin of *E. coli*, DNA sequences encoding the mutant LTs, recombinant expression vectors comprising the DNAs, recombinant microorganisms transformed with the recombinant expression vectors, process for preparing the mutant LTs and pharmaceutical application of the said proteins as immunogenic antigens for vaccination and as adjuvants for antibody production.

### 20 Description of the Prior Art

Enterotoxigenic Escherichia coli("ETEC") causes diarrheal disease in humans and animals due to production of toxin such as heat-labile 25 enterotoxin("LT")(see: Spangler, B.D., Microbiol. Rev., 56:622-647(1992)). LT is a multimeric protein composed of two functionally distinct domains: an enzymatically active A subunit("LTA") of ~30,000daltons with ADP-ribosylating activity, and subunit("LTB") а pentameric В ~11,600daltons that 30 contains GM1 (momosialoganglioside) receptor-binding site(see: Bäckström, M. et al., Microbiol., 24:489-497(1997)). Upon thiol reduction, the A subunit dissociates into two polypeptide chains, i.e., Al(Mr, 23,000 daltons) and A2(Mr, 6,000daltons)(see: Tsuji, 35 T. et al., J. Biol. Chem., 260:8552-8558(1985); Grant C.C.R. et al., Infect. Immun., 62:4270-4278(1994)). subunit, in particular, intoxicates eucaryotic cells by

15

20

25

30

35

catalyzing ADP-ribosylation of Gs, a GTP-binding protein that regulates the levels of the second messenger cAMP(see: Guerrant, R.L. et al., Infect. Immun., 10:320-327(1974); Field, M. et al., N. Engl. J. Med., 321:800-806(1989)). The resulting increase in cAMP level causes secretion of water and electrolytes into the small intestine through interaction with two cAMP-sensitive ion transport mechanisms including (i) NaCl co-transport across the brush border of villous epithelial cells and (ii) electrogenic Na\*-dependent Cl\* secretion by crypt cells(see: Guidry, J.J. et al., Infect. Immun., 65:4943-4950(1997)).

Both the cholera toxin("CT") from Vibrio cholerae and heat-labile enterotoxin from ETEC belong to the most potent mucosal adjuvants and immunogens known to date by oral and other mucosal routes, via which most of antigens are unable induce immune responses (see: Jackson, R.J. et al., Infect. Immun., 61:4272-4279(1993); Takahashi, I. et al., J. Infect. Dis., 173:627-635(1996)). However, their toxicities have precluded their clinical use in humans (see: Douce, G. et al., Proc. Natl. Acad. Sci., USA, 92:1644-1648(1995)). One approach to overcome the problem of toxicity is the generation of genetically detoxified derivatives of LT(see: Lobet, Y. et al., Infect. Immun., 59:2870-2879(1991); Dickson, B.L. and Clements, J.D., Infect. Immun., 63:1617-1623(1995)) and CT(<u>see</u>: Fontana, M.R. et al., Infect. Immun., 63:2356-2360(1995); Yamamoto, S. et al., Proc. Natl. Sci., USA, 94:5267-5272(1997b)) by site-directed mutagenesis of amino acids which are located on the ßstrand that constitutes the 'floor' of NAD-binding cavity.

The most important factor for immunogenicity is shown to be the ability to bind to the receptor on eucaryotic cell(see: Nashar, T.O. et al., Proc. Natl. Acad. Sci., USA, 93:226-230(1996)). In fact, a non-binding mutant of the B subunit of LT was found to be non-immunogenic(see: Guidry, J.J. et al., Infect. Immun., 65:4943-4950(1997)). In addition, another group found that the ADP-ribosylating activity is unnecessary for immunogenicity because nontoxic

10

15

20

25

30

derivatives of LT obtained by site-directed mutagenesis of the A subunit retained the immunological properties of the wild-type LT(see: Pizza, M. et al., J. Exp. Med., 180:2147-2153(1994)).

The attempt to define the role of ADP-ribosylating activity in adjuvanticity of LT has generated conflicting For example, it was reported that a nontoxic derivative of LT(LTE112K) when co-administered with keyhole limpet hemocyanin(KLH) by an oral route in mice, lacked the adjuvant properties, thus suggesting that the adjuvanticity of LT is linked to its ADP-ribosylating activity(see: Lycke, N. et al., Eur. J. Immunol., 22:2277-2281(1992)). However, more recently, the adjuvant activity of the LTE112K was found to be identical to that of the LT holotoxin when delivered with influenza virus surface antigen by intranasal route(<u>see</u>: Verweij, W.R et al., 16:2069-2076(1998)). On the other hand, other investigators showed that another LT derivatives, LTK63, enzymatic activity and toxicity was still able to elicit antibody responses against the co-administered antigen in mice immunized orally, intranasally, or intravaginally (see: Di Tommaso, A. et al., Infect. Immun., 64:974-979(1996); Giuliani, M.M. et al., J. Exp. Med., 187:1123-1132(1998); Marchetti, M. et al., Vaccine, 16:33-37(1998)).

Under the circumstances, the present inventors, based on the findings that detoxified LT derivatives may induce antibody responses, tried to explore an efficient immunogenic antigen and mucosal adjuvant for vaccination, which can be applied for the development of a mucosal vaccine as well as a novel diarrheal vaccine for humans and animals.

### SUMMARY OF THE INVENTION

In accordance with the present invention, the inventors have made an effort to develop detoxified and immunologically active proteins ("mutant LTs") by the site-

directed mutagenesis of heat-labile enterotoxin(LT) of *E. coli*. The inventors cloned full length DNA coding for the LT, mutated the A subunit by site-directed mutagenesis and constructed expression vectors comprising the DNAs of mutated LT and recombinant microorganisms transformed with the recombinant expression vectors. Further, the inventors found that the recombinant mutant LTs can be applied as an active ingredient for diarrheal vaccine and adjuvant for mucosal vaccine.

10

15

20

30

The first object of the invention is, therefore, to provide detoxified and immunologically active proteins which have mutated amino acid sequences of heat-labile enterotoxin of *E. coli*.

The second object of the invention is to provide DNA sequences encoding the said mutant LTs.

The third object of the invention is to provide recombinant expression vectors comprising the said DNAs.

The fourth object of the invention is to provide recombinant microorganisms transformed with the said recombinant expression vectors.

The fifth object of the invention is to provide a process for preparing recombinant mutant LTs from the said microorganisms.

The sixth object of the invention is to provide a diarrheal vaccine comprising an active ingredient of mutant LT.

The seventh object of the invention is to provide a novel use of the mutant LT as an adjuvant for mucosal vaccine.

### BRIEF DESCRIPTION OF THE INVENTION

The above and the other objects and features of the present invention will become apparent from the following description given in conjunction with the accompanying drawings, in which:

10

15

20

25

30

35

Figure 1 is a photograph showing SDS-PAGE of a wildtype and mutant LTs produced isolated from recombinant E. coli. Figure 2 is photographs showing the changes in CHO-K1 cell morphology after treatment wild-type or mutant LTs. Figure is photograph 3 showing ADPribosyltransferase activity of wild-type LT, LTS63Y and 110/112 analyzed by SDS-PAGE followed by autoradiography. Figure is graph showing the amount of intracellular CAMP of CHO cells measured by an enzyme immunoassay system after treatment with wildtype or mutant LTs. Figure 5 is graphs showing anti-LT secretary IgA, serum IgG and IgA antibody responses on intragastric or intranasal immunization. Figure 6 is graphs showing anti-urease secretary IgA, serum IgG and IgA antibody responses on intragastric immunization. Figure 7 is graphs showing anti-urease secretary IgA, serum IgG and IgA antibody responses on intranasal immunization. Figure 8A is a graph showing anti-urease serum IgG subclass antibody responses on

intragastric immunization.

15

20

25

30

35

Figure 8B is a graph showing anti-urease serum IgG subclass antibody responses on intranasal immunization.

### 5 DETAILED DESCRIPTION OF THE INVENTION

The present inventors first isolated a DNA fragment heat-labile enterotoxin(LT) gene enterotoxigenic E. coli K88ac strain of porcine origin and thus cloned into pBluescript KS(-) vector to generate pBlueKS-/rLT for site-directed mutagenesis. Site-directed mutagenesis was performed on single-stranded pBlueKS-/rLT vector. As a result, two vectors that have mutated DNA sequences encoding amino acids of proposed ADPribosyltransferase active center of LT were generated: the first one is 'pBlueKS-/LTS63Y' wherein serine residue at position 63 of LT is substituted with tyrosine; and, the other one is 'pBlueKS-/LT 110/112' wherein glutamic acid residues at positions 110 and 112 are deleted.

Amino acids at positions 58 to 72 are shown to be folded in a  $\beta$ -strand followed by  $\alpha$ -helix, which form the NAD-binding site and amino acids Arg<sup>7</sup>, His<sup>44</sup>, Ser<sup>61</sup> Glu<sup>110</sup> and Glu<sup>112</sup> have been shown to be important for enzymatic activity. Up to date, the modified residues of the LTA subunit are Arg<sup>7</sup>, Ser<sup>63</sup>, Glu<sup>110</sup>, Glu<sup>112</sup> or Ala<sup>72</sup>.

The inventors have substituted Ser<sup>63</sup>, the essential amino acid for NAD-binding and catalytic activity of LT, with Tyr residue having a bulky side chain of phenolic ring. Thus, it was expected to efficiently block NAD-binding in slightly modified LT structure. The inventors have also deleted  $\operatorname{Glu}^{110}$  and  $\operatorname{Glu}^{112}$  in LTA subunit which are important for enzymatic activity, and have studied for immunogenicity to LT itself and immune-eliciting property coadministered antigens. The double deletion mutant of the invention was expected to completely eliminate enzymatic activity of LT.

15

20

25

30

In describing the mutant LTs of the present invention, LTS63Y is employed to mean mutated LT in which serine residue at position 63 is substituted with tyrosine; and, "LT $\Delta$ 110/112" is employed to mean mutated LT in which glutamic acid residues at positions 110 and 112 of LT are deleted, respectively.

pBlueKS-/LTS63Y and pBlueKS-/LT $\Delta$ 110/112, coli transformed into E. Top 10F'. Each transformants thus prepared was designated as 'Escherichia coli Top 10F'-pBlueKS-/LTS63Y' and 'Escherichia coli Top 10F'-pBlueKS-/LT $\Delta$ 110/112' and deposited with the Korean Collection for Type Cultures(KCTC) located at KRIBB #52, Oun-dong, Yusong-gu, Taejon 305-333, Republic of Korea, an international depository authority as accession Nos. KCTC 0648BP and KCTC 0649BP, respectively.

E. coli Top 10F' transformed with either pBlueKS-/LTS63Y or pBlueKS-/LT $\Delta$ 110/112 was grown in LB broth containing 100  $\mu$ g/ml of ampicillin and the mutant LTs were purified from the cultures using immobilized D-galactose column and FPLC Superdex 200 column.

Since the mobilities of mutant LTs were identical to those of the wild-type LT on SDS-PAGE analysis, the molecular weight of the mutant LT subunits were presumed to be identical to those of wild-type LT. These results suggest that mutant LTs consisting of A and B subunits retain the  $AB_5$  conformation similar to wild-type LT.

The ability of mutant LTs to induce morphological changes in cultured Chinese hamster ovary-K1(CHO-K1) cells(ATCC, USA) was tested. The morphological changes in the CHO-K1 cells were used to detect the toxic activity of mutant LTs(see: Grant, C.C.R. et al., Infect. Immun., 62:4270-4278(1994)). As little as 100ng/ml of LT induced longitudinal growth of approximately 90% of the CHO-K1 cells, a response previously shown to be dependent upon adenylate cyclase-induced increases in cAMP(see: Guerrant, R.L. et al., Infect. Immun., 10:320-327(1974)). However, the cells treated with each of the mutant LT at the level

10

15

20

25

35

of  $10\mu g/ml$  showed no morphological changes of the CHO-K1 cells.

ADP-ribosyltransferase activity was determined as the ability to catalyze the transfer of labeled ADP-ribose from [adenylate-32P]NAD to the 41kDa G protein membrane (see: Locht, C. et al., Infect. Immun., 55:2546-2553(1987)). In general, Al subunit of LT is known to catalyze ADP-ribosylation of the membrane-bound substrate G When membrane proteins from CHO-K1 cells were proteins. incubated with wild-type LT in the presence of [adenylateit specifically ADP-ribosylated the Mr 41,000 proteins, which correspond to the  $\alpha$  subunits of the GTP binding Gs protein. In contrast, no ADP-ribosylation of this protein was detected in reaction mixtures incubated with the same amounts of LTS63Y or LT⊿110/112. This result was identical to that of the negative control treated without toxins. Therefore, the substitution of Tyr<sup>63</sup> for Ser<sup>63</sup> or deletion of Glu<sup>110</sup> and Glu<sup>112</sup> in Al subunit appeared to cause changes in structural integrity of NAD binding crevice that may be important for enzymatic activity of LT.

In addition, the levels of cAMP were determined in CHO cells treated with CT, LT, or mutant LTs. The addition of CT or LT caused about 10-fold higher levels of cAMP production than that of untreated cultures. On the other hand, changes in cAMP levels in LTS63Y- or LTA110/112treated cells were barely detectable. These data showed that the presence of A subunit of wild-type LT(accurately subunit) is necessary for an increase in intracellular cAMP concentration and the mutant derivatives, LTS63Y and LT $\Delta$ 110/112, devoid of enzymatic activity, are unable to form cAMP.

The enterotoxicity experiments of mutant LTs examined using a mouse ileal loop test strongly indicate that the mutant LTs possess negligible enterotoxicity in vivo.

In the immunogenic ability of mutant LTs, the mice immunized with LTS63Y or LT $\Delta$ 110/112 contained high and

15

20

25

comparable levels of anti-LT antibodies in sera and fecal extracts compared with those immunized with wild-type LT. Moreover, both of the intragastric and intranasal immunizations using mutant LTs could be an effective method for inducing antibody responses of vaccination.

ability of mutant LTs to act as a mucosal adjuvant was assessed by intragastric immunization in mice. LT 2 110/112 was effective as a mucosal adjuvant intragastric immunization by inducing high levels of mucosal and systemic antibody responses to the coadministered antigens such as H. pylori whole cell lysate or urease. The ability of mutant LTs to function as a mucosal adjuvant was also assessed by intranasal immunization in mice. Intranasal administration of LTS63Y demonstrated the sensitiveness in inducing mucosal immunigenecity and adjuvanticity. Mice immunized intranasally by coadministration of urease antigen and LTS63Y showed strong mucosal and systemic anti-urease responses including urease-specific secretary IgA, serum IgG and IgA antibodies.

The present invention is further illustrated in the following examples, which should not be taken to limit the scope of the invention.

Example 1: Expression and Purification of mutant LTs

Example 1-1: Plasmid construction and mutagenesis

A 1.5Kb BamHI DNA fragment including LT gene from enterotoxigenic E. coli K88ac strain of porcine origin was cloned into pBluescript KS(-) vector(Stratagene, USA). The resulting vector, designated as pBlueKS-/rLT, was used for site-directed mutagenesis.

Site-directed mutagenesis was performed on single-stranded DNA prepared from *E. coli* CJ236 transformed with pBlueKS<sup>-</sup>/rLT according to the method of MutanK kit(Takara

30

35

Biomedicals, Japan). The sequence of oligonucleotides used for the substitution and deletion of amino acids were 5'-ATATGATGACGGATATGTTTCCACTTACCTTAGTTTGAGAAGTGCTCACTTG-3'(SEQ ID NO:1) and 5'-AGGCGTATACAGCCCTCACCCATATCAGGTTTCTGCGTTAGG TGGAATACCAT-3'(SEQ ID NO:2), respectively. As a result, serine residue at position 63 was substituted with tyrosine and glutamic acid residues at positions 110 and 112 were deleted, respectively.

These residues are in proposed ADP-ribosyltransferase active center of LT and their substitutions or deletions have been shown to inactivate ADP-ribosyltransferase activity and enterotoxicity(see: Domenighini, M. et al., Mol. Microbiol., 14:41-50(1994)).

Amino acids at positions 58 to 72 are shown to be folded in a  $\beta\text{-strand}$  followed by  $\alpha\text{-helix,}$  which form the 15 NAD-binding site and amino acids Arg7, His44, Ser61 Glu110 and  $\operatorname{Glu}^{112}$  have been shown to be important for enzymatic activity. Up to date, the modified residues of the LTA subunit are Arg<sup>7</sup>(to Lys), Ser<sup>63</sup>(to Lys), Glu<sup>110</sup>(to Asp), Glu<sup>112</sup>(to Asp or Lys) or Ala<sup>72</sup>(to Arg)(see: Lobet, Y. et al., 20 Infect. Immun., 59:2870-2879(1991); Fontana, M.R. et al., Infect. Immun., 63:2356-2360(1995); Di Tommaso, A. et al., infect. Immun., 64:974-979(1996); Douce, G. et al., Infect. Immun., 63:2821-2828(1997); Marchetti, M.M. et al., Vaccine, 16:33-37(1998); Tsuji, T. et al., FEBS Letters, 292:319-25 321(1991); Lycke, N. et al., Eur. J. Immunol., 22:2277-2281(1992); Verweij, W.R. et al., Vaccine, 2076(1998); Giuliani, M.M. et al., J. Exp. Med., 187:1123-1132(1998)).

The substitution of Ser<sup>63</sup>, the essential amino acid for NAD-binding and catalytic activity of LT, with Tyr residue having a bulky side chain of phenolic ring was expected to efficiently block NAD-binding in slightly modified LT structure. Deletion of Glullo and Glullo, which are located at the putative ADP-ribosyltransferase active center was also expected to completely eliminate enzymatic activity of LT.

25

30

35

The changes of DNA sequences were confirmed using Sequenase Version 2.0 sequencing kit(Amersham Life Science, USA) and each of mutants thus prepared was designated as 'pBlueKS-/LTS63Y' and 'pBlueKS-/LTΔ110/112', respectively.

Example 1-2: Expression and purification of the recombinant mutant LTs

pBluescript KS(-) vectors containing the mutant

LT gene which comprises the 160bp 5'-noncoding region,
1.2kb coding region and 197bp 3'-noncoding region, pBlueKS/LTS63Y and pBlueKS-/LT⊿110/112, were transformed into E.
coli Top 10F' (Invitrogen, USA). Each of the transformants
thus prepared was designated as 'Escherichia coli Top 10F'
pBlueKS-/LTS63Y' and 'Escherichia coli Top 10F'-pBlueKS-/LT

Δ110/112' and deposited with the Korean Collection for
Type Cultures(KCTC) located at KRIBB #52, Oun-dong, Yusonggu, Taejon 305-333, Republic of Korea, an international
depository authority as accession Nos. KCTC 0648BP and KCTC
0649BP, respectively.

E. coli Top 10F' transformed with either LTS63Y or LT  $\Delta$ 110/112 was grown in LB broth containing 100  $\mu$ g/ml of ampicillin and the mutant LTs were purified from the The cells were harvested by centrifugation, resuspended in TEAN buffer(50mM Tris-HCl, pH 7.5, 0.2M NaCl, EDTA and 3mM  $NaN_3$ ), and lysed with microfluidizer (Microfluidics Corporation, USA). The lysates were clarified by centrifugation and then filtered using 0.45 mm membrane (Micro Filtration Systems, Japan) prior to chromatography on an immobilized D-galactose column (Pierce, (see: Uesaka, Y. et al., Microb. Pathog. 76(1974)). Bound proteins were eluted with 0.3M galactose in TEAN buffer. Holotoxin(AB<sub>5</sub>) was separated from the free B-subunit pentamer by size exclusion chromatography using FPLC Superdex 200 column (Pharmacia, Sweden).

The homogeneity of LTS63Y and LT $\triangle$ 110/112 was confirmed by SDS-polyacrylamide gel electrophoresis(see:

15

20

25

30

35

Figure 1). In Figure 1, lanes 1 and 2 represent wild-type LTs; lanes 3 and 4, LTS63Ys; and, lanes 5 and 6, LT $\Delta$ 110/112s, respectively, and the arrow 1 is a position of LT holotoxins and LTB subunit pentamers; arrow 2, LTA subunit; and, arrow 3, LTB subunit monomer. Each lane received 10µg of the proteins and samples in lanes 2, 4 and 6 were heated to 95°C for 5min in the presence of  $\beta$ -mercaptoethanol, while samples in lanes 1, 3 and 5 were loaded without denaturation by heating and adding  $\beta$ -mercaptoethanol. When the purified mutant LTs were analyzed without denaturation, two protein bands appeared: one band with the size of 70-100kDa corresponding to the holotoxin and LTB pentamers; other band with the size of the about corresponding to the LTA subunit. When the purified mutant were boiled for 5min with  $\beta$ -mercaptoethanol, holotoxins were dissociated into two bands of about 30 and 11kDa, corresponding to the A and B subunits of Since the mobilities of mutant LTs were respectively. identical to those of the wild-type LT, the molecular weight of the mutant LT subunits were presumed to be identical to those of wild-type LT.

These results suggest that the innate structure of the A subunit associated with pentameric B subunits of LT is not affected by substitution of tyrosine for  $Ser^{63}$  or deletion of  $Glu^{110}$  and  $Glu^{112}$  residues on NAD-binding pocket. Moreover, it was demonstrated that the binding ability of the B subunit of mutant LTs to  $G_{M1}$  ganglinoside was similar to that of the normal B subunit using a  $G_{M1}$ -ELISA, and mutant LTs were reacted with anti-LT antibody in Western blot analysis. These results imply that mutant LTs retain the AB<sub>5</sub> conformation similar to wild-type LT.

Example 2-1: Cell elongation assay

15

20

25

30

35

PCT/KR99/00555

The ability of mutant LTs to induce morphological in cultured Chinese hamster ovary-K1 (CHO-K1) cells(ATCC, USA) was tested as described previously(see: Guerrant, R.L. et al., Infect. Immun., 10:320-327(1974); Grant, C.C.R. et al., Infect. Immun., 62:4270-4278(1994)). CHO-K1 cells were incubated for 24hr as monolayer cultures in minimal essential medium alpha(MEM-α)(Gibco-BRL, USA) supplemented with 10% fetal bovine serum(FBS) in humidified and 5%  $CO_2$  atmosphere at 37°C. The cells were washed once with Hanks's balanced salt solution(HBSS) (Sigma, USA) and incubated with 0.1% trypsin for 5min. centrifugation, they were washed once, and then resuspended in the growth medium. To each well of a 48-well tissue culture plate, the same numbers of CHO-K1 cells(104  $200\mu l$  volume per well) were allowed to adhere for 4hr prior to the addition of the toxin dilutions and then incubated in humidified and 5% CO, atmosphere at 37°C for 24hr. cells were then washed with phosphate-buffered saline (PBS), fixed with methanol, and stained with 0.04% Trypan Blue Solution(Gibco-BRL, USA). After staining, the cells were washed, air dried, and analyzed for morphological changes by light microscopy.

The morphological changes in the CHO-K1 cells were used to detect the toxic activity of mutant LTs(see: Grant, C.C.R. et al., Infect. Immun., 62:4270-4278(1994)). little as 100ng/ml of wild-type LT induced longitudinal growth of approximately 90% of the CHO-K1 cells, a response previously shown to be dependent upon adenylate cyclaseinduced increases in cAMP(see: Guerrant, R.L. et al., Infect. Immun., 10:320-327(1974)). However, the cells treated with mutant LTs at the level of  $10\mu g/ml$  showed no morphological changes of the CHO-K1 cells (see: Figure 2 and Table 1). In Figure 2, A shows toxin-untreated CHO-K1 cells; B, CHO-K1 cells treated with 100ng/ml of wild-type LT; C, CHO-K1 cells treated with 10µg/ml of LTS63Y; and, D, CHO-K1 cells treated with 10µg/ml of LT 2 110/112, respectively.

35

Table 1. Comparison of biologic and enzymatic activities of wild-type LT, LTS63Y and LT⊿110/112

Toxin assessed	Cell elongationa	Ileal loop test <sup>b</sup>
PBS	10%<	negative
Wild-type LT	90%> at 100ng	positive at 100ng
LTS63Y	10%< at 10μg	negative at 100μg
LT 🛮 110/112	10%< at 10μg	negative at 100µq

a: 10<sup>4</sup> of CHO-K1 cells were cultured with 100ng of wild-type LT or 10µg of each mutant LT for 24hr and a positive toxin effect in the CHO-K1 cells was defined as elongation of >20% of the cells according to published criteria(see: Guerrant, R.L. et al., Infect. Immun. 10:320-327 (1974))

b: The enterotoxicity of mutant LTs was examined using on ileal loop test, where mice were anesthetized, and 100µl of PBS containing 100ng of wild-type LT or 100µg of each mutant LT was injected into an ileal loop. Loops were examined 18hr later and the ratio of fluid to length was defined as positive when the ratio was >40µl/cm(see: Fujita, K. et al., J. Infect. Dis. 125: 647-655 (1972))

### Example 2-2: ADP-ribosyltransferase activity test

20 For the preparation of crude membranes, CHO-K1 cells were maintained in monolayer culture by serial passage in MEM- $\alpha$  medium supplement with 10% FBS(see: Locht C. et al., Immun., 55:2546-2553(1987)). The cells were detached from the flask, washed once in PBS(pH 7.2) and then lysed in a homogenizer (Weaton, USA). The supernatant from these lysates was collected following centrifugation at 1,000xg for 20 min and further centrifuged at 18,000xg for 7 min to obtain microsomal or membrane pellets. The washed membrane pellet was resuspended in 50mM Tris-HCl(pH8.0) at a concentration of 1mg of protein per ml and 30 stored at -70℃ until use.

ADP-ribosyltransferase activity was determined as the ability to catalyze the transfer of labeled ADP-ribose from [adenylate- $^{32}$ P]NAD to the 41kDa G protein in CHO-K1 membrane(see: Locht, C. et al., Infect. Immun., 55:2546-2553(1987)). Reaction mixtures (100 $\mu$ l) containing 32 $\mu$ M [adenylate- $^{32}$ P]NAD(2 $\mu$ Ci) (NEN, USA), 50 mM Tris-HCl(pH 8.0)

10

15

20

25

30

35

containing 10mM thymidine, 100mM ATP, 20mM DTT, 100µM GTP, 50µg of CHO-K1 membrane proteins, and 10µg of wild-type or mutant LTs were incubated at 37% for 30min. Reactions were terminated by the addition of 1ml of ice-cold 50mM Tris-HCl(pH 8.0) and the membranes were sedimented by centrifugation (15,000xg, 7min) at  $4^{\circ}$ C. The membrane pellet was resuspended in ice-cold Tris-HCl(pH 8.0) and sedimented once more by centrifugation before being solubilized in 50µl of electrophoresis sample buffer(67mM Tris-HCl, pH 6.8, 28 SDS, 0.03% bromophenol blue, and 10% containing 5%  $\beta$ -mercaptoethanol. The samples were heated for 5min and then analyzed by SDS-PAGE autoradiography.

In general, the Al subunit of LT is capable binding NAD and catalyzing the ADP-ribosylation membrane-bound GTP-binding regulatory protein Gs, a associated with adenylate cyclase(see: Spangler, B.D., Microbiol. Rev., 56:622-647(1992)). The consequence is a sharp increase in cAMP production resulting in excessive accumulation of salts and water in the intestinal lumen(see: Field, M. et al., N. Engl. J. Med., 322:800-806(1989)). The A subunit of LT is known to catalyze ADPribosylation of the membrane-bound substrate G proteins.

Figure 3 shows ADP-ribosyltransferase activity of LT, LTS63Y and LT $\triangle$ 110/112 analyzed by SDS-PAGE in a 12.5% gel followed by autoradiography. In Figure 3, lane 1 represents a reaction mixture incubated without toxins; lane 2, with 10 $\mu$ g of wild-type LT; lane 3, with 10 $\mu$ g of LTS63Y; and, lane 4, with 10 $\mu$ g of LT $\triangle$ 110/112, respectively, and the arrow denotes the position of the Mr-41,000 band corresponding to the Gs protein.

As shown in Figure 3, when  $50\mu g$  of membrane proteins from CHO-K1 cells were incubated with wild-type LT in the presence of [adenylate- $^{32}P$ ]NAD, it specifically ADP-ribosylated the Mr-41,000 proteins, which correspond to the  $\alpha$  subunits of the GTP binding Gs proteins. In contrast, no ADP-ribosylation of this protein was detected in reaction

mixtures incubated with the same amounts of LTS63Y or LT $\Delta$  110/112. This result was identical to that of the negative control treated without toxins in lane 1. Therefore, the substitution of Tyr $^{63}$  for Ser $^{63}$  or deletion of Glu $^{110}$  and Glu $^{112}$  in A subunit did cause changes in structural integrity of NAD binding crevice that may be important for enzymatic activity of LT.

## Example 2-3: Measurement of intracellular cAMP accumulation

10

15

20

25

30

35

CHO cells (ATCC, USA) were maintained in MEM- $\alpha$  medium FBS in a 24-well supplemented with 10% plate at concentration of 5x10<sup>4</sup> cells per well, grown to near confluency, and incubated in MEM- $\alpha$  containing FBS and 1mM 3-isobutyl-1-methylxanthine (IBMX) for 30min prior addition of toxins(see: Grant, C.C.R. et al., Immun., 62:4270-4278(1994)). Either cholera toxin(CT), cholera toxin B subunit (CTB), trypsin-activated wild-type LT, LTS63Y, or LT⊿110/112 was added to each well and the plates were incubated for 18h. The cells were washed three times with PBS and intracellular cAMP was extracted by adding 200µl of 50mM HCl to each well and placing the plates in -70℃ deep freezer for 20min. cAMP was measured with a Biotrak CAMP enzyme immuno-assay(EIA) system(Amersham Life Science, USA) according to the manufacturer's instructions.

As a result, the levels of cAMP were determined in CHO cells treated with CT, CTB, LT, LTS63Y, or LT 110/112(see: Figure 4). As shown in Figure 4, the addition of CT or LT to reach a concentration of 50ng/ml caused about 10-fold higher levels of cAMP production than those of untreated cultures. On the other hand, cAMP formation in cultures treated with CTB, LTS63Y or LT 110/112 was undetectable even at a concentration as high as 5µg/ml. In Figure 4, 'S' and 'D' denote LTS63Y and LT 110/112, respectively, and the gray bars represent treatments of 500ng of CT or LT and 5µg of CTB, S or D; and, black bars,

50ng of CT or LT and 500ng of CTB, S or D. Results are shown as mean titers and error bars indicate standard deviations from the mean.

These data showed that the presence of wild-type LTA subunit (accurately LTA1 subunit) is necessary for an increase in the intracellular cAMP concentration and the mutant derivatives, LTS63Y and LT 2110/112, devoid of enzymatic activity, are unable to form cAMP.

5

15

. 20

25

30

35

10 Example 2-4: Assessment of toxicity using mouse ileal loops

The enterotoxicity of mutant LTs was examined using a mouse iteal loops test(see: Yamamoto, S. et al., J. Exp. Med., 185:1203-1210(1997)). Groups of mice were anesthetized, and different doses of each toxin were injected into iteal loops(LT, 100ng or lµg per mouse; and, mLT,  $10\mu g$  or  $100\mu g$  per mouse) of individual mice. The mice were sacrificed 18hr after the injection, and the fluid content of the iteal loops was determined; values of more than  $40\mu l/cm$  were considered positive (indicative of toxicity).

One hundred nanogram of wild-type LT induced significant fluid accumulation in small intestine, while no fluid accumulation was observed in the loop treated with thousand-fold higher levels  $(100\mu g)$  of mutant LTs (see: Table 1). These data strongly indicate that the mutant LTs possess negligible enterotoxicity in vivo.

Example 3: Immunological Characterization

Example 3-1: Mucosal immunogenicities of mLTs

Six-week-old female Balb/c mice were purchased from Charles River(Japan). The mucosal immunogenicities of LTS63Y and LT $\Delta$ 110/112 were tested via two immunization routes. Groups of mice were immunized intragastrically with 25 $\mu$ g of LTS63Y or LT $\Delta$ 110/112 four times on days 0, 7,

18

14 and 21 or intranasally with 2μg of LTS63Y or LT⊿110/112 on days 0, 7 and 14(see: Takahashi, I. et al., J Infect. Dis., 173:627-635(1996)). The control groups received PBS alone. The serum and fecal antibody titers to LT were determined using samples prepared on day 7 following the last immunization(see: Figure 5). Figure 5 shows anti-LT secretary IgA, serum IgG and IgA antibody responses on intragastric(white bar) or intranasal(black bar) immunization, where 1 represents PBS treatment; 2, wild-type LT treatment; 3, LTS63Y treatment; 4, LT⊿110/112 treatment.

10

15

20

25

30

35

As shown in Figure 5, the mice immunized with LTS63Y or LT⊿110/112 contained high and comparable levels of anti-LT antibodies in sera and fecal extracts compared with those immunized with wild-type LT. The LTS63Y was slightly more immunogenic than LT⊿110/112 on both intragastric and intranasal administration. On the other hand, titers of anti-LT in the serum or fecal extracts of mice intranasally immunized with wild-type or mutant LTs were slightly higher than those observed in mice intragastrically administered. Intranasal immunization offers several advantages compared with other immunization route: lower doses of proteins are required to induce antibody responses, which means lower cost for vaccine production (see: Yamamoto, S. et al., Proc. Acad. Sci., USA, 94:5267-5272(1997)). administered intranasally, only 6% of the quantity of mutant LT used in intragastric immunization was required to elicit slightly higher levels of secretary IgA responses and this dose also effectively induced systemic IgG and IgA antibody responses. Thus, intranasal immunization using mutant LT could be an effective method for vaccination in humans and animals.

### Example 3-2: Mucosal adjuvanticity of mutant LTs

To test a mucosal adjuvanticity, mice were immunized either intragastrically by 125µg of *H. pylori* urease

15

25

30

35

together with 25µg of LTS63Y or LT $\Delta$ 110/112 on days 0, 7, 14 and 21 or intranasally by 20µg of the same antigen together with 2µg of LTS63Y or LT $\Delta$ 110/112 as an adjuvant on days 0, 7 and 14. Fecal extracts and serum samples were collected on day 7 following the last immunization and the appearance of either mucosal or systemic antibody responses was monitored using ELISA.

LT- and urease-specific antibodies were measured with a G<sub>M1</sub> capture enzyme-linked immunosorbent assay(G<sub>M1</sub>-ELISA) and direct ELISA, respectively, as described Spiegel, S. J. Cell. Biochem., 42:143previously (see: 152(1990); Douce, G. et al., Infect. Immun., 65:2821-2828(1997)). Plates were coated with 150ng of Gmi (Sigma, USA) per well of a 96-well EIA/RIA plate(Costar, USA) for a  $G_{mi}$ -capture ELISA, and then incubated at 37°C for 1hr. Plates were washed three times with PBS containing 0.05% Tween 20("PBST") and blocked with 2.5% skim milk(Difco, USA) in PBST at  $37^{\circ}$ C for 1hr. After washing with PBST three times, 100ng of wild-type LT was added into wells and times with PBST. In case of a direct ELISA, plates were coated with 1µg of urease per well incubated for 1hr at 37°C with horseradish peroxidase (HRP)-conjugated antimouse antibodies specific for mouse immunoglobulinG (IgG) (1:5000), IgA (1:2000) (KPL, USA) or IgG1, IgG2a or IgG2b (1:2000) (Biosource, USA). After washing six times with bound antibody was visualized by addition 3,3'5'5'-tetramethylbenzidine (TMB). The absorbance at 450 nm was determined and ELISA titers were recorded as the highest dilution of serum which gave an absorbance value above the level measeured in preimmune samples.

## Mucosal adjuvanticity of LTS63Y and LT 2110/112 by intragastric immunization

The ability of mutant LTs to act as a mucosal adjuvant was assessed by intragastric immunization in

20

mice(<u>see</u>: Figure 6: 1, PBS treatment; 2, 125 $\mu$ g urease treatment; 3, 125 $\mu$ g urease and 25 $\mu$ g LT treatment; 4, 125 $\mu$ g urease and 25 $\mu$ g LTS63Y treatment; 5, 125 $\mu$ g urease and 25 $\mu$ g LT $\Delta$ 110/112 treatment).

5

10

15

20

25

30

35

As shown in Figure 6, mice immunized intragastrically with H. pyroli urease alone showed no significant levels of to the antigen. Mice immunized coadministration of urease and LT 2110/112 produced high levels of urease antibody responses comparable to those immunized with wild-type LT, while mice immunized coadministration of the antigen and LTS63Y showed significant sero-conversion. Altering the dose of LT⊿ 110/112 to 10 or  $125\mu g$  did not cause any change in the immune response to urease. LTS63Y did not significant adjuvanticity to the urease, even when  $125\mu g$ was coadministered with the urease.

To test the adjuvanticity of LT $\triangle$ 110/112 with other antigens, whole cell lysate of H. pylori was used as an antigen and it was observed that LT $\triangle$ 110/112 also induced antibody responses to whole cell lysate. These data suggest that LT $\triangle$ 110/112 effectively functions as a mucosal adjuvant on intragastric immunization by inducing strong mucosal and systemic antibody responses to coadministered antigens such as H. pylori whole cell lysate or urease.

To further assess the immunologic response induced by LTΔ110/112 following intragastric immunization, IgG subclass responses were determined by ELISA(see: Figure 8A: 1, PBS treatment; 2, 125μg urease treatment; 3, 125μg urease and 25μg LT treatment; 4, 125μg urease and 25μg LTΔ110/112 treatment).

As shown in Figure 8A, urease antibody responses enhanced by  $LT \triangle 110/112$  were largely restricted to IgG1(gray bars), rather than IgG2a(white bars) or IgG2b(black bars) subclass antibody in sera, and similar antibody patterns were observed using wild-type LT as an adjuvant.

Generally, CT elicits adjuvant responses by inducing antigen-specific CD4+ T cells secreting interleukin 4(IL-4), IL-5, IL-6 and IL-10 that correlated directly with serum IgG1 and IgG2b subclass responses in mice orally immunized with protein Ag and CT as adjuvant (see: Marinaro. M. et al., J. Immunol., 155:4621-4629(1995). It was reported that mutant CT(S61F), despite lack of ADP-ribosyltransferase activity with resultant cAMP induction, elicits serum IqG1 IgG2b subclass Ab responses when administrated intranasally(see: Yamamoto, S. et al., Proc. Natl. Acad. 10 USA, 94:5267-5272(1997) or subcutaneously(see: Yamamoto, S. et al., J. Exp. Med., 185:1203-1210(1997). the other hand, oral immunization with LT promotes IgG1, IgG2a and IgG2b, which are supported by a mixed CD4+ Th1and Th2-type responses associated with IFN-y, IL-4, IL-5, 15 IL-6 and IL-10 production(see: Takahashi, I. et al., Infect. Dis., 173:627-635(1996). In the results of present invention, intragastric administration of LT⊿110/112 wild-type LT as an adjuvant induced predominant IgG1 Ab responses as shown in Figure 8A. 20 This result is not consistent with the typical IgG subclass responses induced by wild-type LT and rather resembles the responses induced by CT activating CD4+ Th2-type cells(see: Marinaro, M. et al., J. Immunol., 155:4621-4629(1995). In contrast, LTS63Y 25 induced IgG1, IgG2a and IgG2b antibody responses to H. pylori urease on intranasal immunization. The similar result is also observed by wild-type LT eliciting IgG1, IgG2a and IgG2b subclass responses, which are supported by a mixed CD4+ Th1- and Th2-type response(see: Takahashi, I. et al., J. Infect. Dis., 173:627-635(1996)). Therefore, the mutant LTs of this invention, LTS63Y and LT 2110/112, induced distinct IgG subclass responses, depending on immunization routes.

35 <u>Mucosal adjuvanticity of LTS63Y and LT2110/112 by</u> intranasal immunization

WO 01/19998

10

15

20

25

30

35



The ability of mutant LTs to function as a mucosal adjuvant was also assessed by intranasal immunization in mice. Intranasal administration of wild-type or mutant LTs demonstrated the sensitiveness in inducing mucosal immunigenecity and adjuvanticity(see: Figure 7: 1, PBS treatment; 2, 20µg urease treatment; 3, 20µg urease and 2µg wild-type LT treatment; 4, 20µg urease and 2µg LTS63Y treatment; 5, 20µg urease and 2µg LTD/112 treatment).

Mice immunized intranasally by coadministration of urease antigen and LTS63Y showed high levels of mucosal and systemic anti-urease responses including urease-specific secretary IgA, serum IgG and IgA antibodies, which were equivalent to the responses observed when wild-type LT was used an adjuvant. However, mice immunized coadministration of urease and LT⊿110/112 showed lower level of antibodies to urease, similar to those levels induced by urease alone. When 0.2µg of LTS63Y was used, antibody responses to urease in both fecal extracts and sera were lower by a factor of approximately 10, but when 6μg of LTS63Y was used, no increase in antibody responses to urease was induced.

In addition, as shown in Figure 8B, LTS63Y induced IgG1(gray bars), IgG2a(black bars) and IgG2b(white bars) subclass antibody responses to *H. pylori* urease, and these results were quite different from those IgG subclass responses(predominant IgG1) resulted from the intragastric immunization with the mutant LT\(\triangle 110/112\)(see: Figure 8B: 1, PBS treatment; 2, 20\(\triangle g\) urease treatment; 3, 20\(\triangle g\) urease and 2\(\triangle g\) LTS63Y treatment; 5, 20\(\triangle g\) urease and 2\(\triangle g\) LT\(\triangle 110/112\) treatment). Therefore, the mutant LTs of this invention, LTS63Y and LT\(\triangle 110/112\) induced district IgG subclass responses, depending on immunization routes.

As described above, the mutant LTs of this invention have shown different abilities to act as mucosal adjuvant according to the route of administration. This suggests that different mutant forms of LT may require different

15

20

25

30

immunization routes for adequate adjuvanticity, and that specific immunization route may elicit specific mechanisms of up-regulation of immune responses, independent of cAMP levels. Thus, selection of the route of immunization may be critical for determining the mucosal adjuvant activity of mutant LTs. Mucosal vaccines delivered into the nasal tract provide several advantages. For example, lower doses of antigen and adjuvant are required to induce effective antibody responses when compared to intragastric immunization, which can decrease the cost of vaccination.

As clearly illustrated and demonstrated as above, the present invention provides detoxified and immunologically active proteins (mutant LTs). The mutant LTs was expressed recombinant expression vectors, pBlueKS-/LTS63Y and pBlueKS-/LT 110/112 that contain mutated DNA sequences encoding amino acids in the ADP-ribosyltransferase active In contrast to wild-type LT, both of the LTS63Y and LT⊿110/112 did not induce any toxic activities. of the mutants elicited high and comparable levels of anti-LT antibodies when delivered either intragastrically or intranasally, inducing systemic and local responses serum and fecal extracts. Thus, they might be useful for the development of a novel diarrheal vaccine in human and In addition, the antibody production ability using LTS63Y and LT 2110/112 as adjuvants against H. pylori urease may be effective for prevention and treatment of various diseases. The adjuvant activity of these mutants might be very useful to develop an effective mucosal vaccine component.



BUDAPEST TREATY ON THE INTERNATIONAL RECOGNITION OF THE DEPOSIT OF MICROORGANISMS FOR THE PURPOSE OF PATENT PROCEDURE

### INTERNATIONAL FORM

### RECEIPT IN THE CASE OF AN ORIGINAL DEPOSIT

issued pursuant to Rule 7.1

TO: Mogam Biotechnology Research Institute
#341, Pojung-ri, Kuusung myust, Yongin si, Kyonggi-do 4/0-910,
Republic of Korea

### I. IDENTIFICATION OF THE MICROORGANISM

Identification reference given by the DEPOSITOR:

Escherichia coli Top10F'-pBlueKS'/LTS63Y Accession number given by the INTERNATIONAL DEPOSITARY AUTHORITY:

**KCTC 0648BP** 

### D. SCIENTIFIC DESCRIPTION AND/OR PROPOSED TAXONOMIC DESIGNATION

The microorganism identified under I above was accompanied by:

[x] a scientific description

[ ] a proposed taxonomic designation (Mark with a cross where applicable)

### III. RECEIPT AND ACCEPTANCE

This International Depositary Authority accepts the microorganism identified under I above, which was received by it on Jul 27 1999.

### N. RECEIPT OF REQUEST FOR CONVERSION

The microorganism identified under I above was received by this international Depositary Authority on and a request to convert the original deposit to a deposit under the Budapest Treaty was received by it on

### V. INTERNATIONAL DEPOSITARY AUTHORITY

Name: Korean Collection for Type Cultures

Address: Korea Research Institute of

Bioscience and Biotechnology

(KRIBB)

#52, Oun-dong, Yusong-ku,

Taejon 305-333, Republic of Korea Signature(s) of person(s) having the power to represent the International Depositary Authority of authorized official(s):

BAE, Kyung Sook, Director Date: Aug 02 1999



### BUDAPEST TREATY ON THE INTERNATIONAL RECOGNITION OF THE DEPOSIT OF MICROORGANISMS FOR THE PURPOSE OF PATENT PROCEDURE

### INTERNATIONAL FORM

### RECEIPT IN THE CASE OF AN ORIGINAL DEPOSIT

issued pursuant to Rule 7.1

TO: Mogam Biotechnology Research Institute
#341, Pojung-ri, Koosung-myun, Yongin-si, Kyonggi-do 449-910,
Republic of Korea

### I. IDENTIFICATION OF THE MICROORGANISM

Identification reference given by the DEPOSITOR:

Escherichia coli Top10F'-pBlueKS'/LT4110/112. Accession number given by the INTERNATIONAL DEPOSITARY AUTHORITY:

KCTC 0649BP

### II. SCIENTIFIC DESCRIPTION AND/OR PROPOSED TAXONOMIC DESIGNATION

The microorganism identified under I above was accompanied by:

[ x ] a scientific description

[ ] a proposed taxonomic designation (Mark with a cross where applicable)

### III. RECEIPT AND ACCEPTANCE

This International Depositary Authority accepts the microorganism identified under I above, which was received by it on Jul 27 1999.

### IV. RECEIPT OF REQUEST FOR CONVERSION

The microorganism identified under I above was received by this International Depositary Authority on and a request to convert the original deposit to a deposit under the Budapest Treaty was received by it on

### V. INTERNATIONAL DEPOSITARY AUTHORITY

Name: Korean Collection for Type Cultures

Address: Korea Research Institute of Bioscience and Biotechnology

(KRIBB)

#52. Oun-dong, Yusong-ku.

Taejon 305-333, Republic of Korea Signature(s) of person(s) having the power to represent the International Depositary Authority of authorized official(s):

BAE, Kyung Sook, Director Date: Aug 02 1999



WO 01/19998

- 1. A detoxified and immunologically active protein having an amino acid sequence of heat-labile enterotoxin of *E. coli*, in which position 63 or positions 110 and 112 is mutated in a site-directed manner.
- 2. A detoxified and immunologically active protein having an amino acid sequence of heat-labile enterotoxin of *E. coli*(SEQ ID NO:3), in which serine residue at position 63 is substituted with tyrosine.
- 3. A DNA sequence (SEQ ID NO:4) encoding the detoxified and immunologically active protein of claim 2.
  - 4. A recombinant expression vector pBlueKS-/LTS63Y which comprises the DNA of claim 3.
- 5. Escherichia coli Top10F'-pBlueKS-/LTS63Y(KCTC 0648BP) transformed with the recombinant expression vector pBlueKS-/LTS63Y of claim 4.
- 6. A detoxified and immunologically active protein having an amino acid sequence of heat-labile enterotoxin of E. coli(SEQ ID NO:5), in which glutamic acid residues at positions 110 and 112 are deleted.
- 7. A DNA sequence(SEQ ID NO:6) encoding the detoxified and immunologically active protein of claim 6.
  - 8. A recombinant expression vector pBlueKS-/LT $\Delta$  110/112 which comprises the DNA of claim 7.
- 9. Escherichia coli Top10F'-pBlueKS-/LT⊿110/112(KCTC 0649BP) transformed with the recombinant expression vector pBlueKS-/LT⊿110/112 of claim 8.

- A process for preparing the detoxified and immunologically active protein(mutant LT) of claim 1 which comprises the steps of culturing a recombinant microorganism transformed with an expression comprising a DNA encoding the mutant LT and isolating the recombinant mutant LT from the culture.
- 11. The process for preparing the detoxified and immunologically active protein of claim 10, wherein the recombinant microorganism is *Escherichia coli* Top10F'-pBlueKS-/LTS63Y(KCTC 0648BP) or *Escherichia coli* Top10F'-pBlueKS-/LT⊿110/112(KCTC 0649BP).
- 15 12. A diarrheal vaccine comprising an active ingredient of the detoxified and immunologically active protein of claim 1 and pharmaceutically acceptable carrier.
- 13. A mucosal adjuvant comprising an active ingredient of the detoxified and immunologically active protein of claim 1.

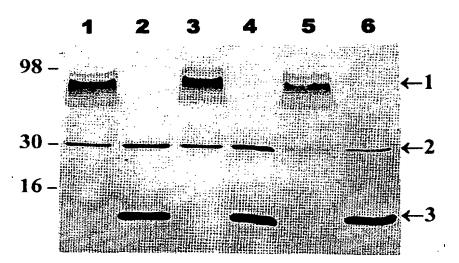


Fig. 1

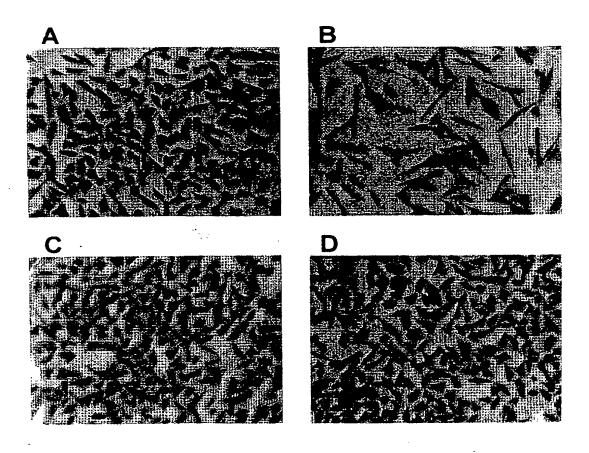


Fig. 2

1 2 3 4

98 -

30 –

16 –

Fig. 3

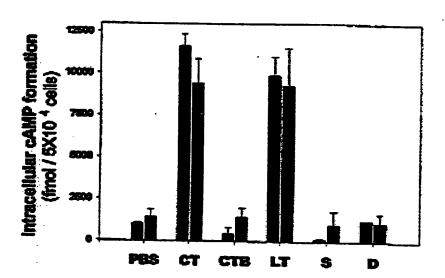
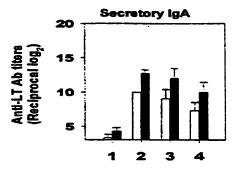
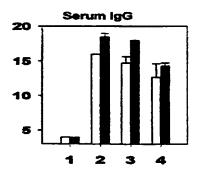


Fig. 4





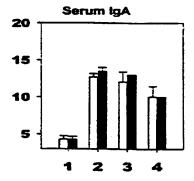


Fig. 5

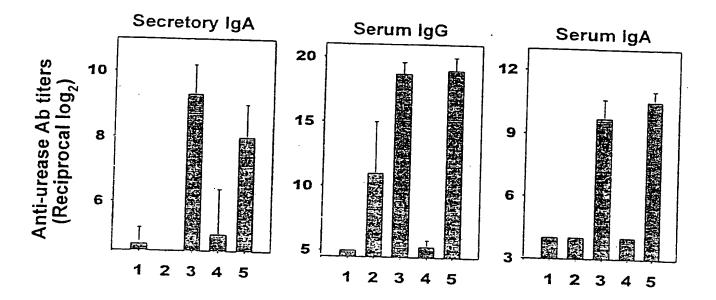


Fig. 6



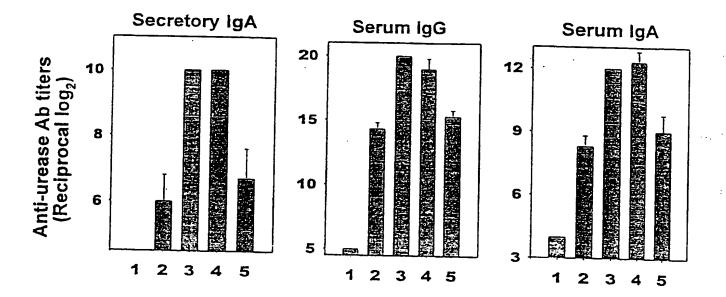


Fig. 7

8/9

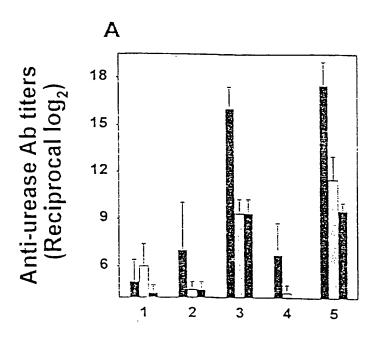


Fig. 8A

9/9

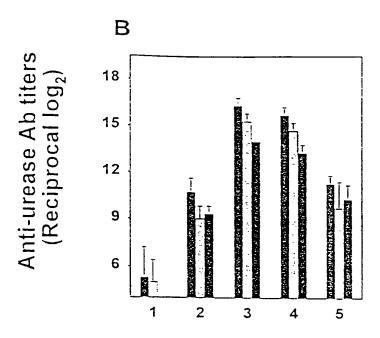


Fig. 8B





## SEQUENCE LISTING

5	<110>	MOGAM BIOTECHNOLOGY RESEARCH INSTITUTE et al.
	<120>	NOVEL DETOXIFITED MUTANTS OF Escherichia coli HEAT-LABILE ENTEROT OXIN
10	<160>	6
10	<170>	KOPATIN 1.5
	<210>	1
	<211>	52
15	<212>	DNA
	<213>	Artificial Sequence
	<220>	
	<223>	single-stranded nucleotide
20		
	<400>	1
	atatgatga	e ggatatgttt ccacttacct tagtttgaga agtgctcact tg 52
25	<210>	2
	<b>&lt;211&gt;</b>	53
	<212>	DNA
	<213>	Artificial Sequence
30	<220>	
	<223>	single-stranded oligonucleotide
	<400>	2
	aggcgtata	c ageceteace catateaggt ttetgegtta ggtggaatac cat 53
35		
	<210>	3





	<211>	382					
	<212>	PRT					
	<213>	Escherichia co	li		•		
5	<400>	3					
	Met Lys	Asn Ile Thr Phe I	le Phe Phe Ile	Leu Leu Ala S	er Pro Leu		
	1	5		10		15	
	Tvr Ala A	Asn Gly Asp Arg	Leu Tvr Ara A	la Asn Ser Are	Pro Pro Asn		
10	- y	20	La Iyi Aig A	25	30 s		
		20		23	50		
	Glu Ile L	ys Arg Ser Gly G	ily Leu Met Pro	o Arg Gly His	Asn Glu Tyr		
		35	40		45		
15	Phe Asp	Arg Gly Thr Gln	Met Asn Ile A	sn Leu Tyr As <sub>l</sub>	His Ala Arg		
	50		55		60		
		Gin Thr Gly Phe			Val Ser Thr		
20	65		70 ·	75			80
20	Twelens	Ser Leu Arg Ser	Alo Uio I ou Al	a Clu Cla So-	llo I au Cas		
	Tyr Dea .	Sei Leu Aig Sei 7	Ala HIS LEU AI	a Gly Gin Ser 90	ile Leu Ser	95	
		63		90		93	
	Gly Tyr S	Ser Thr Tyr Tyr I	le Tvr Val Ile A	la Thr Ala Pro	Asn Met		
25		100		05	110		
	Phe Asn	Val Asn Asp Val	Leu Gly Val T	ут Ser Pro His	Pro Tyr Glu		
		115	120		125		
		,					
30	Gln Glu	Val Ser Ala Leu (	Gly Gly Ile Pro	Tyr Ser Gin II	e Tyr Gly		
	130		135		140		
		Arg Val Asn Phe			His Arg Asn		
26	145	1	150	155			160
35	Ara Glui	Tur Ara Ara Ara	Tractice Assa	on Lau A 11-	Alo Des Al-		
	AIG CILL	Tyr Arg Asp Arg	Tyr Tyr Arg A	sn Leu Asn He 170	AIA PTO AIA	175	



	Glu Asp Gly Tyr Arg I	Leu Ala Gly Phe Pro	Pro Asp His Gln Ala	Ггр
	180	18:	5	190
5	Arg Glu Glu Pro Trp I 195	le His His Ala Pro G 200	ln Gly Cys Gly Asn S 205	er .
10	Ser Arg Thr Ile Thr Gi	ly Asp Thr Cys Asn o	Glu Glu Thr Gln Asn I 220	Leu
	Ser Thr Ile Tyr Leu Ar 225	rg Glu Tyr Gln Ser L 230	ys Val Lys Arg Gln Ile 235	240
15	Phe Ser Asp Tyr Gln S	_	yr Asn Arg Ile Arg As 250	sp 255
	Glu Leu Met Asn Lys 260	Val Lys Phe Tyr Val 26.		Leu 270
20	Ser Ser Leu Cys Ala H	lis Gly Ala Pro Gln :	Ser Ile Thr Glu Leu Cy	/s
	275	280	285	
	Ser Glu Tyr His Asn T	hr Gln Ile Tyr Thr II	e Asn Asp Lys Ile Leu	l
25	290	295	300	
	Ser Tyr Thr Glu Ser M	let Ala Gly Lys Arg	Glu Met Val Ile Ile Th	r
	305	310	315	320
	Phe Lys Ser Gly Ala T	hr Phe Gln Val Glu	Val Pro Gly Ser Gln H	is
30	32	.5	330	335
	lle Asp Ser Gln Lys Ly	ys Ala Ile Glu Arg M	let Lys Asp Thr Leu A	<b>r</b> g
	340	34	5	350
35	Ile Thr Tyr Leu Thr G	lu Thr Lys Ile Asp L	ys Leu Cys Val Trp As	sn
	355	360	365	





# Asn Lys Thr Pro Asn Ser Ile Ala Ala Ile Ser Met Glu Asn 370 375 380

5	<210>	4	
	<211>	1514	
	<212>	DNA	
	<213>	Escherichia coli	
10	<400>	4	٠
	ggatccgtg	c actettett tategettea etacacattt tateetegea tggatgtttt	60
	ataaaaaac	a tgattgacat catgttgcat ataggttaaa caaaacaagt ggcgttatct	120
15	ttttccggat	tgictictig tatgatatat aagtitteet egaatgaaaa atataaettt	180
	cattttttt at	ttttattag categeeatt atatgeaaat ggegacagat tatacegtge	240
20	tgactctaga	a cccccagatg aaataaaacg ttccggaggt cttatgccca gagggcataa	300
	tgagtacttc	gatagaggaa ctcaaatgaa tattaatctt tatgatcacg cgagaggaac	360
	acaaaccgg	ge ttigteagat atgatgaegg atatgttice acttacetta gttigagaag	420
25	tgctcactta	gcaggacagt ctatattatc aggatattcc acttactata tatatgttat	480
	agcgacago	ca ccaaatatgt ttaatgttaa tgatgtatta ggcgtataca gccctcaccc	540
30	atatgaacag	g gaggtttetg egttaggtgg aataceatat teteagatat atggatggta	600
50	tcgtgttaat	tttggtgtga ttgatgaacg attacatcgt aacagggaat atagagaccg	660
	gtattacaga	a aatctgaata tagctccggc agaggatggt tacagattag caggtttccc	720
35	accggatca	c caagettgga gagaagaace etggatteat catgeaceae aaggttgtgg	g 780
	aaattcatca	agaacaatca caggtgatac ttgtaatgag gagacccaga atctgagcac	840





	aatatatete a	agggaatatc aatc	aaaagt taagagg	cag atattttcag act	tatcagtc	900	
5	agaggttgac	atatataaca gaa	ttcggga tgaatta	tga ataaagtaaa at	tttatgtt	960	
5	ttatttacgg o	egttactate etetet	tatgt gcacacgga	g etecteagte tatta	acagaa	1020	
	ctatgttcgg	aatatcacaa caca	acaaata tatacga	taa atgacaagat ac	tatcatat	1080	
10	acggaatcga	a tggcaggcaa aa	agagaaatg gttat	catta catttaagag o	eggegeaaca	1140	
	tttcaggtcg	aagtcccggg ca	gtcaacat atagac	tece aaaaaaaage	cattgaaagg	1200	
1 =	atgaaggaca	a cattaagaat cad	catatetg acegag	acca aaattgataa a	ittatgtgta	1260	
15	tggaataata	aaacccccaa ttc	aattgeg geaatea	igta tggaaaacta g	tttgcttta	1320	
	aaagcatgto	taatgctagg aac	cctatata acaacta	ctg tacttatact aat	gagcctt	1380	
20	atgctgcatt	tgaaaaggeg gta	gaggatg caatac	egat cettaaactg t	aacactata	1440	
	acagetteca	ctacagggag ct	gttatage aaacag	aaaa aactaagcta	ggctggaggg	1500	
25	gcaagcttgg	g atcc					1514
	<210>	5					
	<211>	380					
20	<212>	PRT					
30	<213>	Escherichia c	oli				
	<400>	5					
			Ile Phe Phe Ile	: Leu Leu Ala Se	er Pro Leu		
25	1	5		10		15	
35	Tim A1- 4	on Clay A a co	- I T A	Al- A 6 A	D P- 4	_	
	iyr Ala A		g Leu 1yt Arg	Ala Asp Ser Arg		-	
		20		25	3(	U	



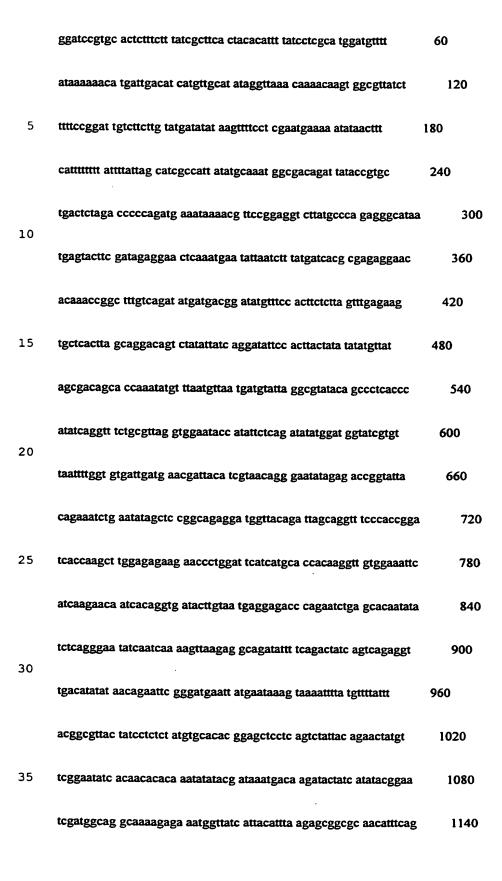
	Glu Ile Lys Arg Ser Gl	y Gly Leu Met P	o Arg Gly His	Asn Glu Tyr	
	35	40		45	
5	Phe Asp Arg Gly Thr	Gln Met Asn Ile A	Asn Leu Tyr As	p His Ala Arg	
	50	55		60	
	Gly Thr Gln Thr Gly F	Phe Val Arg Tyr A	sp Asp Gly Ty	r Val Ser Thr	
10	65	70	75		80
10	Ser Leu Ser Leu Arg S	ser Ala His Leu A	la Gly Gln Ser	Ile Leu Ser	
	8	5	90		95
	Gly Tyr Ser Thr Tyr Ty	yr Ile Tyr Val Ile A	Ala Thr Ala Pro	Asn Met	
15	100		105	110	
	Phe Asn Val Asn Asp	Val Leu Gly Val 1	yr Ser Pro His	Pro Tyr Gln	
	115	120		125	
20	Val Ser Ala Leu Gly G	ily Ile Pro Tyr Sei	Gln Ile Tyr Gl	y Trp Tyr	
	130	135		140	
	Arg Val Asn Phe Gly	Val Ile Asp Glu A	rg Leu His Arg	Asn Arg Glu	
25	145	150	155		160
	Tyr Arg Asp Arg Tyr	Гуг Arg Asn Leu	Asn Ile Ala Pro	Ala Glu Asp	
	16	5	170		175
	Gly Tyr Arg Leu Ala (	Gly Phe Pro Pro A	sp His Gln Ala	Trp Arg Glu	
30	180		185	190	
	Glu Pro Trp Ile His Hi		Cys Gly Asn	Ser Ser Arg	
	195	200		205	
35	Thr Ile Thr Gly Asp T		lu Thr Gln Ası	Leu Ser Thr	
	210	215		220	





	lle Tyr L	eu Arg Glu Tyr (	Gln Ser Lys Val I	ys Arg Gln Ile P	he Ser	
	225	2	230	235		240
	Asp Tyr	Gln Ser Glu Val	Asp Ile Tyr Asn	Arg Ile Arg Asp	Glu Leu	
5		245		250	2:	55
	Met Asn	Lys Val Lys Phe	Тут Val Leu Phe	Thr Ala Leu Le	u Ser Ser	
		260	20	55	270	
10	Leu Cys	Ala His Gly Ala	Pro Gln Ser Ile	Thr Glu Leu Cys	Ser Glu	
		275	280		285	
	Tyr His A	Asn Thr Gln Ile T	ут Thr Ile Asn A	Asp Lys Ile Leu S	er Tyr	
15	290		295	300	1	
	Thr Glu	Ser Met Ala Gly	Lys Arg Glu Me	t Val Ile Ile Thr	Phe Lys	
	305	3	310	315		320
	Ser Gly A	Ala Thr Phe Gln	Val Glu Val Pro	Gly Ser Gln His	Ile Asp	
20		325		330	33	35
	Ser Gln I	Lys Lys Ala Ile G	lu Arg Met Lys	Asp Thr Leu Arg	g Ile Thr	
		340	34	15	350	
25	Tyr Leu	Thr Glu Thr Lys	lle Asp Lys Leu	Cys Val Trp Asn	Asn Lys	
		355	360		365	
	Thr Pro A	Asn Ser Ile Ala A	la Ile Ser Met G	lu Asn		
30	370		375	380		
	<210>	6				
	<211>	1508				
	<212>	DNA				
35	<213>	Escherichia co	li			
	<400>	6				









	gtcgaagtcc cgggcagtca acatatagac tcccaaaaaa aagccattga aaggatgaag	1200	
5	gacacattaa gaatcacata tctgaccgag accaaaattg ataaattatg tgtatggaat	1260	
J	aataaaaccc ccaattcaat tgcggcaatc agtatggaaa actagtttgc tttaaaagca	1320	
	tgtctaatgc taggaaccta tataacaact actgtactta tactaatgag ccttatgctg	1380	
10	catttgaaaa ggcggtagag gatgcaatac cgatccttaa actgtaacac tataacagct	1440	
	tccactacag ggagctgtta tagcaaacag aaaaaactaa gctaggctgg aggggcaagc	1500	
	ttggatcc		1508





International application No. PCT/KR 99/00555

#### CLASSIFICATION OF SUBJECT MATTER

IPC<sup>7</sup>: C 12 N 15/31, 15/09, 1/21; A 61 K 39/108 // (C 12 N 1/21, C 12 R 1:19)

According to International Patent Classification (IPC) or to both national classification and IPC

#### B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC<sup>7</sup>: C 12 N 15/31, 15/09, 1/21; A 61 K 39/108

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)

WPI, CAS, STN-registry

### C. DOCUMENTS CONSIDERED TO BE RELEVANT

Further documents are listed in the continuation of Box C.

Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
EP 0620850 B1 (CHIRON S.P.A.) 3 March 1999 (03.03.99) page 2, lines 17-25; claims 1,3,5,11,19.	1,10,12,13
WO 96/06627 A1 (THE ADMINISTRATORS OF THE TULANE EDUCATIONAL FUND) 7 March 1996 (07.03.96) amended claims 1,4-12,14-17,19,23-25.	1-13
WO 98/28005 A1 (SMITHKLINE BEECHAM BIOLOGICALS S.A.) 2 July 1998 (02.07.98) claims 1,2,6,7,9.	1,13
	[
	EP 0620850 B1 (CHIRON S.P.A.) 3 March 1999 (03.03.99) page 2, lines 17-25; claims 1,3,5,11,19.  WO 96/06627 A1 (THE ADMINISTRATORS OF THE TULANE EDUCATIONAL FUND) 7 March 1996 (07.03.96) amended claims 1,4-12,14-17,19,23-25.  WO 98/28005 A1 (SMITHKLINE BEECHAM BIOLOGICALS S.A.) 2 July 1998 (02.07.98)

"A" document defining the general state of the art which is not considered to be of particular relevance  "E" earlier application or patent but published on or after the international filing date  "L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)  "O" document referring to an oral disclosure, use, exhibition or other means  "P" document published prior to the international filing date but later than the priority date claimed	"I alter document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention "X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone "Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art "&" document member of the same patent family
Date of the actual completion of the international search	Date of mailing of the international search report
17 May 2000 (17.05.2000)	2 August 2000 (02.08.2000)
Name and mailing address of the ISA/AT Austrian Patent Office Kohlmarkt 8-10; A-1014 Vienna Facsimile No. 1/53424/535	Authorized officer  Mosser  Telephone No. 1/53424/437

See patent family annex.

Form PCT/ISA/210 (second sheet) (July 1998)





### INTERNATIONAL SEARCH REPORT

Information on patent family members

International application No. PCT/KR 99/00555

Patent document cited in search report					Patent family member(s)		Publication date
EP	A1	620850	26-10-1994	AT	E	177145	15-03-199
EP	81	620850	03-03-1999	AU	A1	33476/93	28-07-199
				DE	C0	69228563	08-04-199
				DE	T2	69228563	29-07-199
				EP	Al	869181	07-10-199
				ES	т3	2127808	01-05-199
				GR	Т3	3029556	30-06-199
				ΙT	AO	913513	31-12-19
				IT	A	1253009	10-07-199
				JP	T2	7506240	13-07-199
				MX	A1	9207685	31-05-199
				SG	A1	48217	17-04-199
				WO	A1	9313202	08-07-199
WO	A1	9606627	07-03-1996	UA	A1	32337/95	22-03-19
				AU	B2	709779	09-09-19
				BR	A	9508633	30-09-19
				CA	AA	2198586	07-03-19
				CN	A	1164191	05-11-19
				CZ	A3	9700562	15-10-19
				EP	Al	777490	11-06-19
				EP	A4	777490	18-11-19
				FI	A0	970799	26-02-19
				FI	A	970799	24-04-19
				HU	A2	77869	28-09-19
			•	JР	Т2	10505059	19-05-19
				NO	A0	970993	04-03-19
				МО	A	970993	25-04-19
				NZ	Α	291262	25-02-19
				PL	A1	318930	21-07-19
				ບຣ	A	6019982	01-02-20
				ZA	A	9506412	11-03-19
WO	A1	9828005	02-07-1998	AU	A1	57639/98	17-07-19
				EP	A1	948352	13-10-19
				GB	A0	9626864	12-02-19
		·		ZA	A	9711565	27-08-19